FIGURE 1A

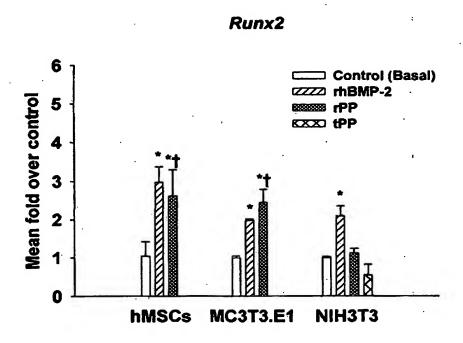


FIGURE 1B

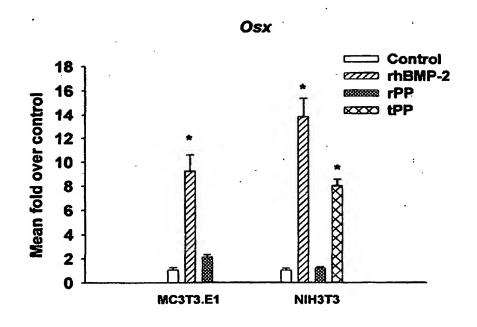


FIGURE 1C

rhBMP-2 induction of Osx in hMSC

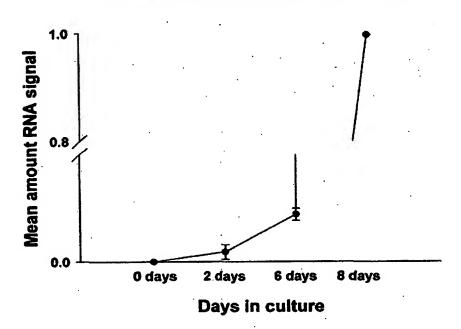


FIGURE 1D

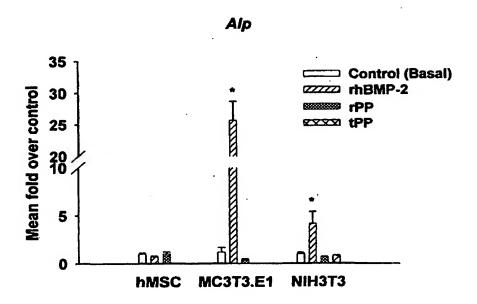


FIGURE 1E

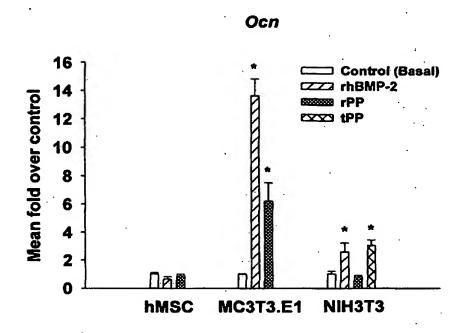


FIGURE 1F

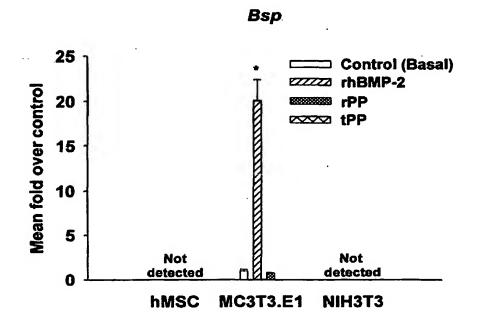


FIGURE 2A

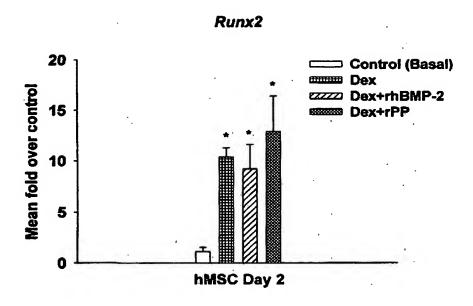


FIGURE 2B

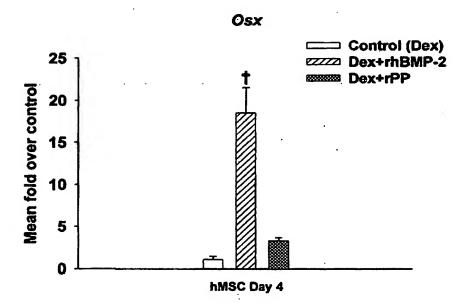


FIGURE 2C

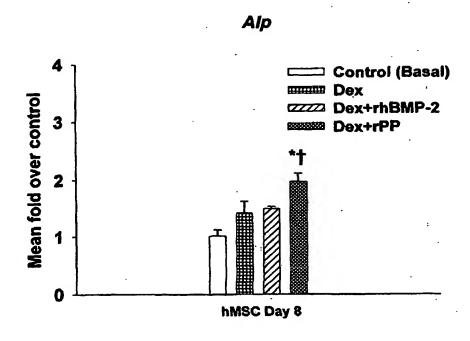


FIGURE 2D

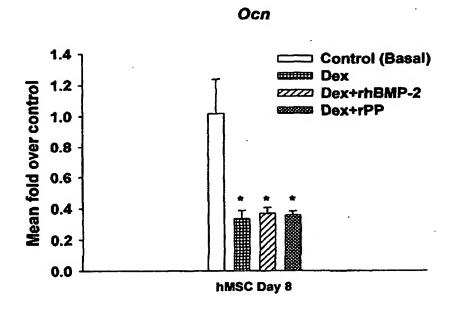


FIGURE 2E

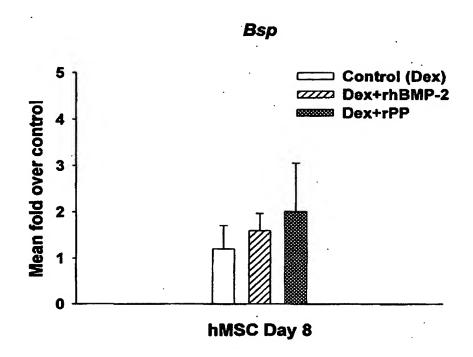


FIGURE 3

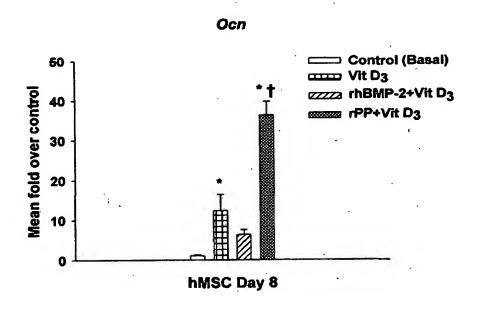


FIGURE 4A

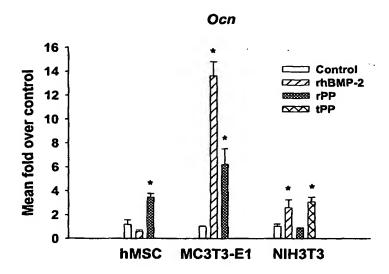


FIGURE 4B

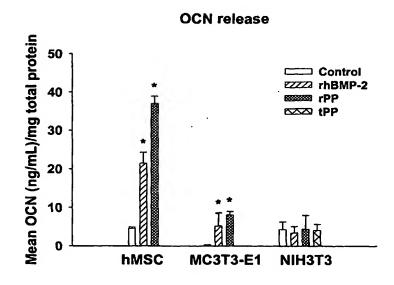


FIGURE 5



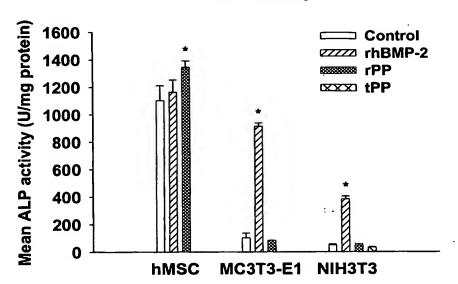


FIGURE 6

FIGURE 6A

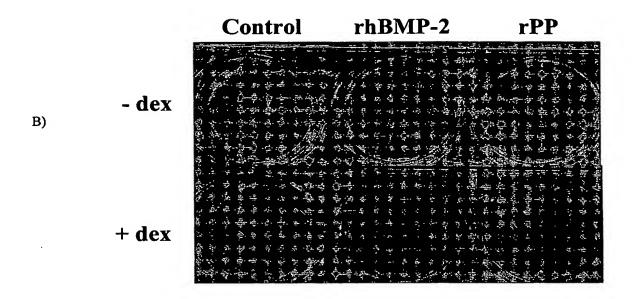


FIGURE 6B

Quantitative alizarin red stain of hMSC

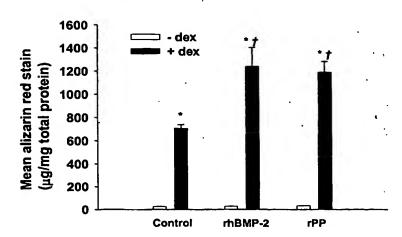
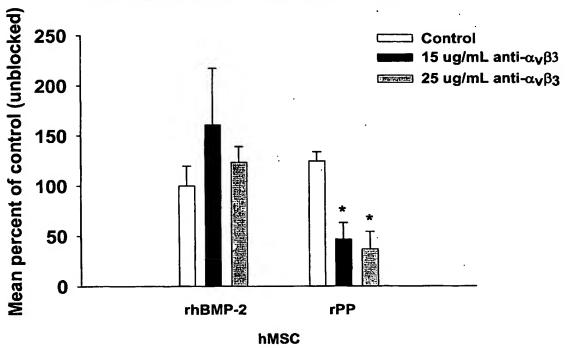


FIGURE 7





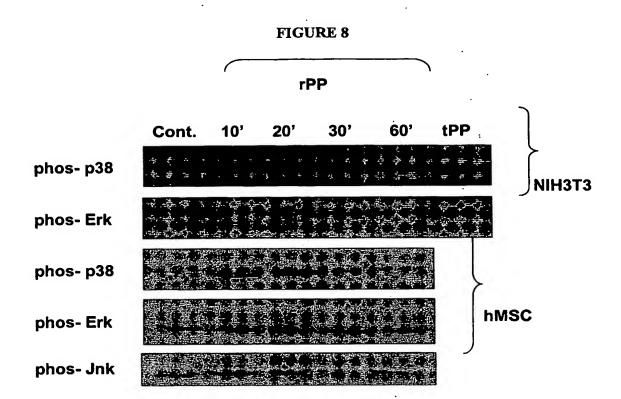


FIGURE 9

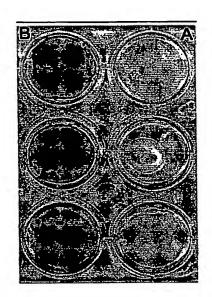
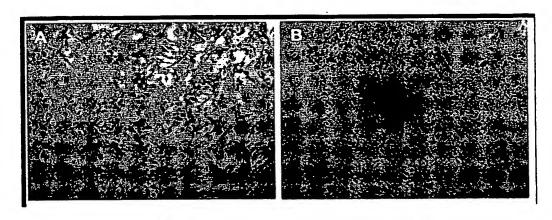


FIGURE 10



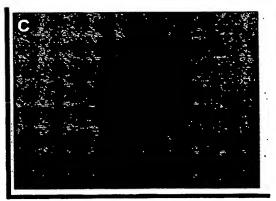


FIGURE 11

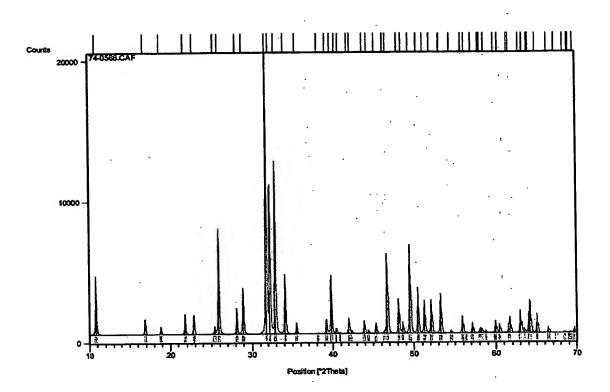


FIGURE 12

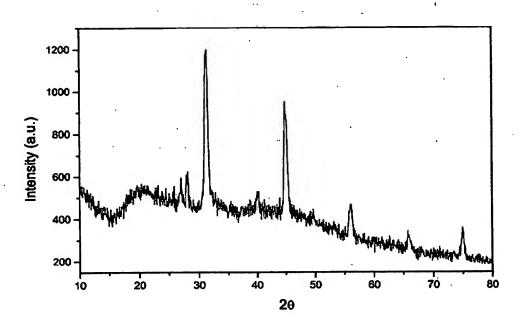
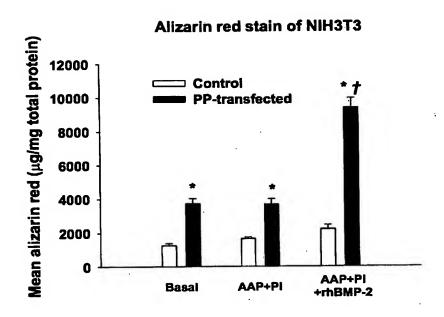


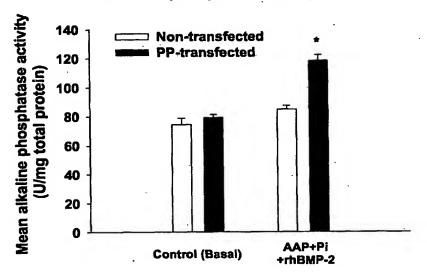
FIGURE 13



* Significant from control, p<0.05
† Significant from PP-transfected, AAP+Pi, p<0.05

FIGURE 14

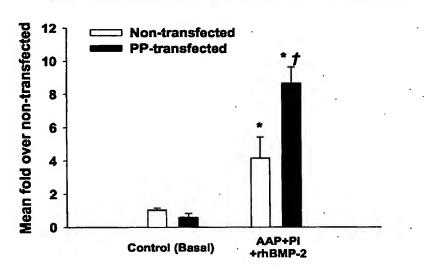
Alkaline phosphtase activity in NIH3T3



^{*} Significant from PP-transfected (Control) and non-transfected (AAP+PI+rhBMP-2), p<0.05

FIGURE 15

Alkaline phosphatase gene expression in NIH3T3



* Significant from Control (Basal), p<0.05
† Significant from non-transfected (AAP+PI+rhBMP-2), p<0.05

FIGURE 16

SEQ ID NO:1

Amino acid sequence of full length Mus musculus PP protein (Exon 5 of DMP3)

SEQ ID NO: 2

Amino acid sequence of a portion of Exon 5 of the Mus musculus PP protein

SEQ ID NO: 3

Nucleotide sequence encoding full length Mus musculus PP protein (Exon 5 of DMP3)

21/37

SEQ ID NO:4 Nucleotide sequence encoding a portion of Exon 5 of the *Mus musculus* PP protein

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SEQ ID NO: 5 Amino acid sequence of *Mus musculus* DMP3 protein GenBank Accession No. AAD42781

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301
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421
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481
541
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721	sassdsssss	dssdsssssd	ssdssdssds	sdssessdss	nssdssdssd	ssdssdssds
781	sdssdssdss	nssdssdssd	ssdssdssns	sdssdssdss	dssdssdssd	ssdssdssds
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SEQ ID NO: 6 Nucleotide sequence of *Mus musculus* DMP3 GenBank Accession No. AP 135799

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421 ttcaaatctg ctattacaac caagttagga agaggaaggc aattcctgag gaaagtggca
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SEQ ID NO:7 Amino acid sequence of human phosphophoryn GenBank Accession No. NP 055023

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SEQ ID NO:8 Nucleotide sequence of human phosphophoryn GenBank Accession No. NM_014208

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2521	actostaces	acagcaacga	tastageage	agcagcgaca	gcagtgatag acagcagtga	cagcaacagc
2581	agreatagea	gtaacagcag	teatagtagt	gatageageg	acagcagtga	tagcgacagc
2641	agcaacagca	acaacaataa	caatagtagt	gacagcagcg	atagcagcaa	cagcagtgat
2701	agtagtgata	gtagtgatag	Cagugacage	agigatageg	acagcagcaa gcaacagcag	tagaagtgac
2761	gatagtagta	acagegacag	Cagegatage	agugacagca	gcaacagcag	tgacagcagt
2821	adcaacadca	atastaata	cagcaatgaa	aycaycaata	gcagtgacag gcagtgacag	cagtgatagc
2881	agcaacagca	atastaacea	trassreset	aycaycaaca	gcagtgacag acaacagcaa	cagtgatagc
2941	agcagcaaca	grantagray	cantraters	aataytagtg	gtaatagtag	tagcagtgac
3001	aatagcaata	acarcarcas	cagcagtage	aytyacayca	gcaatagtag	Lgacagcagc
3061	dacadcadca	acageageaa	tagcagtgac	agragragata	gcaatagcag	cgacagcagt
3121	agtgatagea	acaacaacaa	tratarcart	aycaytyata	acagcagtga	tagegacage
3181	agtagtgaca	deadeageag	cartracare	accaycaycy	gtgacagcag	cagcagtaat
3241	dacaccacta	acaccaccag	tagegacage	aycyatayca	gugacagcag	cgatagcagt
227	gucugcagig	acaycaycaa	caycaytyac	aycaytgaca	gcagcgacag	cagtgatagc

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SEQ ID NO: 9

Amino acid sequence of BMP-2

GenBank Accession Number: NP 001191

```
1 mvagtrclla lllpqvllgg aaglvpelgr rkfaaassgr pssqpsdevl sefelrllsm 61 fglkqrptps rdavvppyml dlyrrhsgqp gspapdhrle raasrantvr sfhheeslee 121 lpetsgkttr rfffnlssip teefitsael qvfreqmqda lgnnssfhhr iniyeiikpa 181 tanskfpvtr lldtrlvnqn asrwesfdvt pavmrwtaqg hanhgfvvev ahleekqgvs 241 krhvrisrsl hqdehswsqi rpllvtfghd gkghplhkre krqakhkqrk rlkssckrhp 301 lyvdfsdvgw ndwivappgy hafychgecp fpladhlnst nhaivqtlvn svnskipkac 361 cvptelsais mlyldenekv vlknyqdmvv egcgcr
```

SEQ ID NO: 10 Nucleotide sequence of human BMP-2 GenBank Accession number NM 001200

```
1 ggggacttct tgaacttgca gggagaataa cttgcgcacc ccactttgcg ccggtgcctt
 61 tgccccagcg gagcctgctt cgccatctcc gagccccacc gcccctccac tcctcggcct
121 tgcccgacac tgagacgctg ttcccagcgt gaaaagagag actgcgcggc cggcacccgg
181 gagaaggagg aggcaaagaa aaggaacgga cattcggtcc ttgcgccagg tcctttgacc
241 agagtttttc catgtggacg ctctttcaat ggacgtgtcc ccgcgtgctt cttagacgga
301 ctgcggtctc ctaaaggtcg accatggtgg ccgggacccg ctgtcttcta gcgttgctgc
361 ttccccaggt cctcctgggc ggcgcggctg gcctcgttcc ggagctgggc cgcaggaagt
421 tegeggegge gtegteggge egeceteat eccagecete tgacgaggte etgagegagt
481 tegagttgcg getgeteage atgtteggee tgaaacagag acceaecece ageagggaeg
541 ccqtqqtqcc cccctacatg ctagacctgt atcgcaggca ctcaggtcag ccgggctcac
601 ccgccccaga ccaccggttg gagagggcag ccagccgagc caacactgtg cgcagcttcc
661 accatgaaga atctttggaa gaactaccag aaacgagtgg gaaaacaacc cggagattct
721 tetttaattt aagttetate eecaeggagg agtttateae etcageagag etteaggttt
781 tecgagaaca gatgeaagat getttaggaa acaatageag tttecateac egaattaata
 841 tttatgaaat cataaaacct gcaacagcca actcgaaatt ccccgtgacc agacttttgg
901 acaccaqqtt ggtqaatcaq aatgcaagca ggtgggaaag ttttgatgtc accccgctg
961 tgatgcqqtg gactgcacag ggacacgcca accatggatt cgtggtggaa gtggcccact
1021 tggaggagaa acaaggtgtc tccaagagac atgttaggat aagcaggtct ttgcaccaag
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1081	atgaacacag	ctggtcacag	ataaggccat	tgctagtaac	ttttggccat	gatggaaaag
		ccacaaaaga				
1201	agtccagctg	taagagacac	cctttgtacg	tggacttcag	tgacgtgggg	tggaatgact
1261	ggattgtggc	tccccgggg	tatcacgcct	tttactgcca	cggagaatgc	ccttttcctc
1321	tggctgatca	tctgaactcc	actaatcatg	ccattgttca	gacgttggtc	aactctgtta
1381	actctaagat	tcctaaggca	tgctgtgtcc	cgacagaact	cagtgctatc	tcgatgctgt
		gaatgaaaag				gtggagggtt
1501	gtgggtgtcg	ctagtacagc	aaaattaaat	acataaatat	atatata	

SEQ ID NO: 11

GGATGGAGCTGTATCATCCTCTTCTTGGTAGCAACAGCTACA

SEQ ID NO: 12

CTAATGTCGACATGGAGAGTGGCAGCCGTGGAGA

SEQ ID NO: 13

GCATTCTAGATTAAAGCACCCGCCATTCAAATCG

FIGURE 17

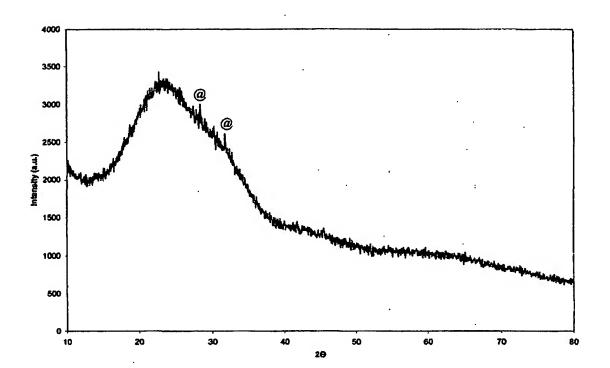


FIGURE 18

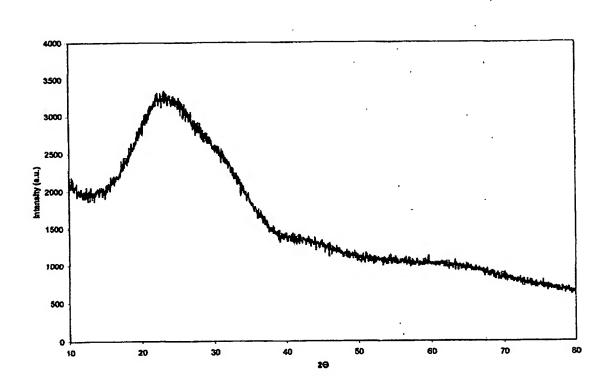


FIGURE 19

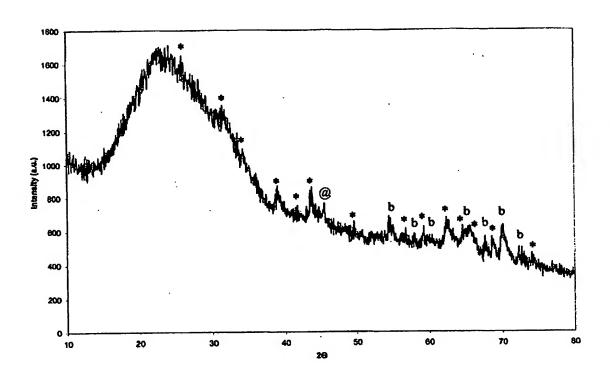


FIGURE 20

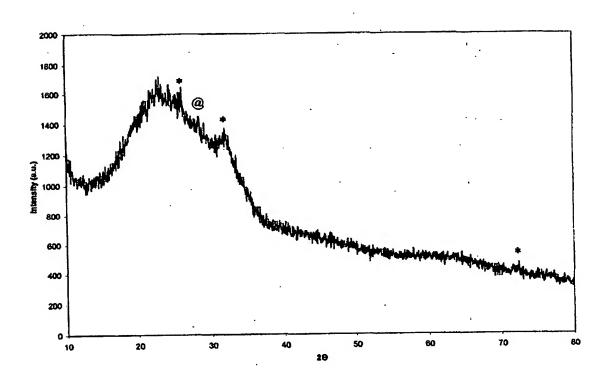
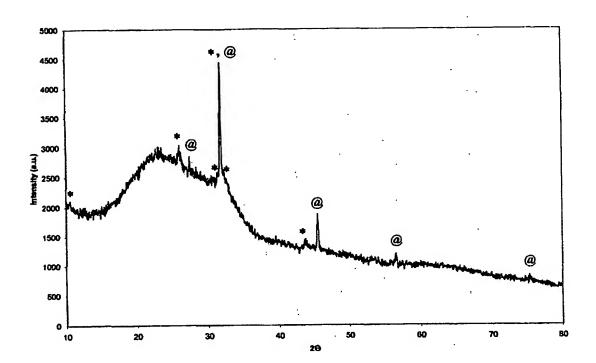


FIGURE 21



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FIGURE 22



FIGURE 23

